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Minimum
Maximum
                                                                                                                                                                                                                                                                                                                                     Result
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Title:
Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                           score greater than or equal to the score of the result bein and is derived by analysis of the total score distribution.
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DB seq length: 2000000000
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118.5
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74.5
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73
73
72.5
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1: /SIDS1/gcgdata/geneseq/geneseqp/Aa1980.DAT:*

2: SIDS1/gcgdata/geneseq/geneseqp/Aa1981.DAT:*

3: /SIDS1/gcgdata/geneseq/geneseqp/Aa1982.DAT:*

4: /SIDS1/gcgdata/geneseq/geneseqp/Aa1983.DAT:*

5: /SIDS1/gcgdata/geneseq/geneseqp/Aa1985.DAT:*

6: /SIDS1/gcgdata/geneseq/geneseqp/Aa1985.DAT:*

7: /SIDS1/gcgdata/geneseq/geneseqp/Aa1985.DAT:*

8: /SIDS1/gcgdata/geneseq/geneseqp/Aa1989.DAT:*

9: /SIDS1/gcgdata/geneseq/geneseqp/Aa1989.DAT:*

10: /SIDS1/gcgdata/geneseq/geneseqp/Aa1999.DAT:*

11: /SIDS1/gcgdata/geneseq/geneseqp/Aa1991.DAT:*

13: /SIDS1/gcgdata/geneseq/geneseqp/Aa1993.DAT:*

14: /SIDS1/gcgdata/geneseq/geneseqp/Aa1993.DAT:*

16: /SIDS1/gcgdata/geneseq/geneseqp/Aa1993.DAT:*

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18: /SIDS1/gcgdata/geneseq/geneseqp/Aa1999.DAT:*

19: /SIDS1/gcgdata/geneseq/geneseqp/Aa1999.DAT:*

20: /SIDS1/gcgdata/geneseq/geneseqp/Aa1999.DAT:*

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21: /SIDS1/gcgdata/geneseq/geneseqp/Aa1999.DAT:*

22: /SIDS1/gcgdata/geneseq/geneseqp/Aa1999.DAT:*
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SIDS1/gcgdata/geneseq/yeneseqp/AA1984.DAT:*

SIDS1/gcgdata/geneseq/geneseqp/AA1985.DAT:*

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Human androgen rec
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Macadamia integrif
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JP;						·									rumo	g's	rer rer	Amino acid sequenc	Protein regulating	GST-HD fusion prot	Mouse brain CNG-1	Amino acid sequenc Piq p105 zona pell	Human ZC1 protein.	Human metastasis-a	Spinocerebellar at Human SCAl protein	Antimicrobial maiz	70K Ul snRNA bindi	70K 7	Peptide seq ID No:	nog	uman andro	liver cance	A human trichohyal

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Sequence
                                                   animals.
                                                                       The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and man
                                                                                                                                                        Claim 1; Page 43-45;
                                                                                                                                                                                                       Novel anti-microbial protein from e.g. Macadamia integrifolia -useful for controlling microbial infestations of plants or mammals
                                                                                                                                                                                                                                                                                              N-PSDB; V42316.
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625
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AA;
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1..28
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Query Match

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Score

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                                                                                         Query Match 94.1%;
Best Local Similarity 92.1%;
Matches 58; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity 93.7 Matches 59; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Key
Peptide
                                                                                                                                                 Sequence
                                                                                                                                                                         animals.
                                                                                                                                                                                 The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and map {\bf map}
                                                                                                                                                                                                                      Claim 1;
                                                                                                                                                                                                                                          Novel anti-microbial protein from e.g. Macadamia integrifolia - useful for controlling microbial infestations of plants or mammals
                                                                                                                                                                                                                                                                            WPI; 1998-377279/32.
N-PSDB; V42310.
                                                                                                                                                                                                                                                                                                             Bower NI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Macadamia integrifolia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       antimicrobial protein; infestation; control
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Macadamia integrifolia antimicrobial protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           W62828;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                W62828 standard; Protein; 666 AA
                                                                                                                                                                                                                                                                                                                                                        20-DEC-1996;
                                                                                                                                                                                                                                                                                                                                                                              22-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                    02-JUL-1998
                                                                                                                                                                                                                                                                                                                                                                                                                         WO9827805-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   27-OCT-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                 (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY
                                         186 krdpqqreyedcrrrceqqeprqqhqcqlrcreqqrqhgrggdmmnpqrggsgryeegee 245
 246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     61 KQS 63
|||
205 kqs 207
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                      61 KQS 63
                                                       1 KRDPQOREYEDCRRRCEQQEPRQQYQCQRRCREQQQRQHGRGGDLINPQRGGSGRYEEGEE 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 KRDPQQREYEDCRRRCEQQEPRQQYQCQRRCREQQRQHGRGGDLINPQRGGSGRYEEGEE 60
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eqs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  krdpqqreyedcrrhceqqeprlqyqcqrrcqeqqrqhgrggdlmnpqrggsgryeegee 204
 248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       207
                                                                                                                                                                                                                   Page 34-36;
                                                                                                                                                                                                                                                                                                            Goulter KC,
                                                                                                                                                  666 AA;
                                                                                                                                                                                                                                                                                                                                                       96AU-0004275
                                                                                                                                                                                                                                                                                                                                                                              97WO-AU00874.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "signal peptide"
29..666
                                                                                                                                                                                                                                                                                                                                                                                                                                                /note=
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
1..28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         93.78;
                                                                                                                                                                                                                   96pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                              "mature protein"
                                                                                                                                                                                                                                                                                                            Green JL,
                                                                                                                                                                                                                    English.
                                                                                        Score 332; DB 19;
Pred. No. 4.5e-30;
4; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pred. No. 2.5e-30; 2; Mismatches 2;
                                                                                                                                                                                                                                                                                                            Manners JM,
                                                                                                                                                                                                                                                                                                             Marcus
                                                                                                               Length 666;
                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                   and mammalian
                                                                                                                                                                                                                                                                                                             JP;
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                                                                                        Gaps
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RESULT
R20181
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                                                                                                                                                                                            Db
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                                                                                                                                                                                                                 Query Match
Best Local Similarity 41.2
11-JUN-1990;
              07-JUN-1991;
                                                        Theobroma cacao
                                                                     Cocoa; flavour; vicilin; seed storage protein.
                                                                                   Sequence encoded by 67 kD T. cacao protein cDNA
                                                                                                                                                                                                                                                           Sequence
                            26-DEC-1991
                                                                                                  16-APR-1992
                                                                                                                            R20181 standard; Protein; 566
                                                                                                                                                                                                                                                                              The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and mammalian
                                                                                                                                                                                                                                                                                                                 Novel anti-microbial protein from e.g. Macadamia useful for controlling microbial infestations of
                                                                                                                                                                                                                                                                                                                                        WPI; 1998-377279/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       antimicrobial protein; infestation; control.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Theobroma cacao antimicrobial protein.
                                                                                                                                                                                                                                                                                                     Claim 1;
                                                                                                                                                                                                                                                                                                                                                                   (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                               22-DEC-1997;
                                                                                                                                                                                                                                                                                                                                                                                                             02-JUL-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Theobroma cacao
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 27-OCT-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 W62831;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              W62831 standard; Protein; 525
                                                                                                                                                                                                                                                                                                                                                                                  20-DEC-1996;
                                                                                                                                                                                            35
                                                                                                                                                                               55
                                                                                                                                                                92 cqeqqqgq 99
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                                                                                                                                                                         YEEGEEKQ 62
                                                                                                                                                                                                                                                                                                   Page 47-49; 96pp;
                                                                                                                                                                                                                                                                                                                                                     Goulter KC,
                                                                                                                                                                                                                                                           525
                                                                                                 (first entry)
                                                                                                                                                                                                                                                            AA;
90GB-0013016
              91WO-GB00914
                                                                                                                                                                                                                                                                                                                                                                                  96AU-0004275
                                                                                                                                                                                                                                                                                                                                                                                              97WO-AU00874.
                                                                                                                                                                                                                       33.6%; Score 118.5; DB 19;
41.2%; Pred. No. 6.3e-06;
ative 14; Mismatches 17;
                                                                                                                                                                                                                                                                                                                                                     Green JL,
                                                                                                                                                                                                                                                                                                   English.
                                                                                                                              AΑ
                                                                                                                                                                                                                                                                                                                                                     Manners JM,
                                                                                                                                                                                                                                                                                                                                                     Marcus JP;
                                                                                                                                                                                                                                                                                                                 integrifolia -
plants or mammals
                                                                                                                                                                                                                        Indels
                                                                                                                                                                                                                                     Length
                                                                                                                                                                                                                                      525;
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W62832
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Best Local S
Matches 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The inventors claim a 67 kD and 31 kD T. cacao protein, and fragments, and encoding DNAs. The 47 kD and 31 kD proteins are derived from the 67 kD precursor. T. cacao protein cDNA was detected in a cDNA library prepared from immature cocca beans RNA using a probe based on the AA sequence of a CNBr peptide common to the 47 kD and 31 kD polypeptides. Homology searches revealed close homologies between the 67 kD polypeptide and the vicilins, which are seed storage proteins.
The sequence is that of an antimicrobial protein which can
                                    Claim 1;
                                                                Novel anti-microbial protein from e.g. Macadamia integrifolia useful for controlling microbial infestations of plants or mamm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Recombinant cocca proteins - are responsible for flavour in cocca beans and produced in large quantities using yeast and bacterial expression vectors
                                                                                                                                                       Bower NI,
                                                                                                                                                                                                                                                                                         02-JUL-1998
                                                                                                                                                                                                                                                                                                                        W09827805-A1
                                                                                                                                                                                                                                                                                                                                                         Gossypium hirsutum
                                                                                                                                                                                                                                                                                                                                                                                        antimicrobial protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             W62832;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           W62832 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 4; Fig 2; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1992-024418/03.
N-PSDB; Q20377.
                                                                                                                                                                                      (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY
                                                                                                                                                                                                                         20-DEC-1996;
                                                                                                                                                                                                                                                        22-DEC-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           92 cqeqqqgq
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         35 erdprq-qyeqcqrrceseateeregeqceqrcereykeqqrqqee--elqrqyqqcqgr 91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            55 YEEGEEKQ 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 KRDPQQREYEDCRRRCEQQ--EPRQQYQCQRRC----REQQRQHGRGGDLINPQRGGSGR 54 :|||:|| :|| :|| :|| :|| :|| :||
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           l Similarity 41.2
28; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ME,
                                  Page 49-51;
                                                                                                                                                     Goulter KC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               566 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99
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                               96pp; English
                                                                                                                                                                                                                                                                                                                                                                                          infestation; control
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                                                                                                                                                     Green JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           590
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Pred. No. 6.8e-06;
4; Mismatches 17;
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                                                                                                                                                       Manners JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ashton
                                                                                                                                                     Marcus
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 566;
                                                                                                                                                       JP;
                                                                     mammals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           9;
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Db
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   RESULT
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W62837
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Best Local Similarity 37.0'
Matches 34; Conservative
                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 22; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  be used animals.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; Page 60-62; 96pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel anti-microbial protein from e.g. Macadamia integrifolia - useful for controlling microbial infestations of plants or mammals % \left( 1\right) =\left\{ 1\right\} 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1998-377279/32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bower NI,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hordeum vulgare
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         antimicrobial protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                               |: ||| ::
102 rhgegeree 110
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                                                                                                                                    54 RYEEGEEKQ 62
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                                                                                                                                                                                                                             qqcvqrcrqerprysharcvqecrddqqqhgrheqeeeqgrgrgwhgegereeehgrgrg 101
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     637
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  590 AA;
                                                                                                                                                                                                                                                                                                                                                           29.2%; Score 103; DB 19; ilarity 31.9%; Pred. No. 0.00044; Conservative 11; Mismatches 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ΑĄ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            96AU-0004275
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             infestation; control
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Green JL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AΑ
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                                                                                                                                                                                                                                                                                                                                                                                                                               Length 637;
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P93109
ID P93109
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 21.1%; Score 74.5; I Best Local Similarity 29.5%; Pred. No. 0.71 Matches 18; Conservative 10; Mismatches
14-APR-1988;
                                                             13-APR-1989;
                                                                                                                                                                                                                                                                                                                            Human androgen receptor; ployclonal antibody; cancer.
                                                                                                                                                                                                                                                                                                                                                                                             Human androgen receptor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                          19-MAR-1990 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               P93109 standard; protein; 919 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The sequence is that of an antimicrobial protein which can be used to control microbial infestations in plants and mammalian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; Page 58-60; 96pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel anti-microbial protein from e.g. Macadamia integrifolia - useful for controlling microbial infestations of plants or mammals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antimicrobial
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                                                                                                                                                                                              WO8909791-A.
                                                                                                                                                                                                                                                            Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1998-377279/32.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (RETR-) COOP RES CENT TROPICAL PLANT PATHOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20-DEC-1996;
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                                                                                                                              19-OCT-1989
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             62 Q 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       40 cvrrcedrpwhqrprcleqcreeerekrqersrheaddrsge-----gssedereqek 92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12 CRRRCEQQEPRQQYQCQRRCREQQRQ------HGRGGDLINPQRGGSGRYEEGEEK 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            593 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein;
                                                             89WO-US01548
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88US-0182646
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Best Local
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           Human androgen receptor (W14783) binds testosterone and, acting at the transcriptional level, regulates the growth of normal prostatic cells. Antisense oligonucleotides (see also T63200,
                                                                                  Oligonucleotide(s) antisense to human androge FGF genes - used to inhibit gene expression, benign prostatic hyperplasia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                anurogen receptor protein (AR) is used to produce mono- or poly-clonal antibodies. These are used for the detection and quantification of AR in the presence of endogenous androgen, as androgen will not interfere with binding. They may be used in assays to determine and quantify cellular distribution of AR in tumour tissue, and are esp. useful for evaluating prostate cancers to determine responsiveness to androgen withdrawal therapy.
  prostatic cells.
T63404-05) based
                                                                Disclosure;
                                                                                                                                                    WPI; 1997-202879/18
                                                                                                                                                                                                                                                                                                                                                                                                                                                W14783
                                                                                                                                                                                                      (WORC-) WORCESTER FOUND BIOMEDICAL RES
                                                                                                                                                                                                                               20-SEP-1995;
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                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                     Androgen receptor; acidic fibroblast growth factor; aFGF; antisense; benign prostatic hyperplasia; prostate cancer;
                                                                                                                                                                                                                                                                                                                                                                                             Androgen receptor
                                                                                                                                                                                                                                                                                                                                                                                                                      22-JUN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        W14783 standard;
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                                                             51pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 73;
Pred. No.
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                                                                                                  human androgen receptor and acidic e expression, for the treatment of
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    useful for transforming
and subsequent antibody |

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(see also
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RESULT PT 8914 FINE PT 8914 FIN
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                                                                                                          androgen receptor, as activation domains, and as a tool for screening for compounds which affect androgen-independent activation of the AR. The peptides, when used in combination with androgen deprivation, effectively limit androgen mediated disease progression. These diseases include cancer, benign prostatic hyperplasia, hirusutism, androgenic alopecia, acne, breast cancer, Kennedy disease, and especially prostate cancer. The peptides and nucleic acids encoding them, are especially used for the
                                                                                                                                                                                                                                                                                                        This sequence represents the human androgen receptor (AR) amino acid sequence. The invention relates to a fragment of the AR corresponding to amino acids 234-391 (see Y78913). The fragment mediates androgen-independent activation of the AR. The androgen receptor acts as a transcription factor, regulating the expression of certain androgen-responsive genes. Interaction of the AR with the protein kinase A signal transduction pathway involves interaction with the androgen independent region. The AR fragment and peptides derived from it can be used as agents for inhibiting androgen independent activation of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    such
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Disclosure; Page 7; 32pp;
                                                                                         treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         androgen-independent activation of
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (UYBR-) UNIV BRITISH COLUMBIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-JUN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23-MAY-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            compounds and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bruchovsky N,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      919 AA;
                                                           androgen-mediated diseases, especially rived of androgen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 tor; AR; androgen-independent activation; inhibitor;
prostatic hyperplasia; hirusutism; androgenic alopecia;
ancer; Kennedy disease; prostate cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98US-0091871.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    cancer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            for treatment of androgen-mediated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gout PW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 919
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 73;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        amino acid sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Snoek R,
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Sequence

919 AA;

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RESULT
Y30795
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Best Local Similarity
Tatches 19; Conserv?
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                                                                        Query Match
Best Local S
Matches 19
                                                                                                                                        The protein is used to raise antibodies affinity purification. The 98 kD product starts at the first Me starts from the second.
                                                                                                                      Sequence
                                                                                                                                                                                     Claim 8; Fig 3; 60pp; English.
                                                                                                                                                                                                         purification.
                                                                                                                                                                                                                 New DNA encoding
to bind DNA, and
                                                                                                                                                                                                                                                WPI; 1989-309501/42
N-PSDB; N91577.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-FEB-1990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             P90996 standard;
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                                                                         l Similarity
19; Conserv
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                                                                                                                       919 AA;
                                                                         Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      receptor DNA clone.
                                                                                                                                                                                                                                                                                                                                                                                                        /product=98 kD
185..919
                                                                                                                                                                                                                                                                                                                                                                                                                          /*tag=
                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
1..919
                                                                                                                                                                                                                                                                                                                                                                                       /product=79 kD polypeptide
                                                                                                                                                                                                                  new androgen receptor and TR2 polypeptide(s) - ab derived antibodies, useful for receptor assay and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20.7%;
                                                                        20.5%; Score 72.5; D
32.8%; Pred. No. 1.9;
tive 12; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              919
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 73; DB 2
Pred. No. 1.6;
15; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A
                                                                                                                                                                                                                                                                                                                                                                                                                   polypeptide
                                                                                                                                                  first Met codon; the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 21; Length 919;
                                                                                                                                                                   for receptor
                                                                                          DB
                                                                                          10;
                                                                         22;
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                                                                                                                                                79 kD product
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                                                                        Gaps
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RESULT
Y33504
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Best Local Similarity
""" hes 22; Conserve
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                                                                                                                                                                                                                                                                                                                                                                                                Proapoptotic; dependence domain; p75NTR; androgen receptor; DCC; huntingtin polypeptide; Machado-Joseph disease; SCA1; SCA2; SCA6; atrophin-1; cell death; apoptosis; Huntington's disease; head trauma; Alzheimer's disease; Kennedy's disease; spinocerebellar ataxia; stroke; dentatorubropallidoluysian atrophy; cell proliferation; cell survival; neoplastic; malignant; autoimmune; fibrotic.
                                                                                                                                                                                                                                                                                                                                                                                                                                The present sequence represents a human trichohyalin (TRHY) protein. The protein is found in terminally differentiating epidermal tissue, and is involved in forming the structural architecture of such tissue. The trichohyalin protein is useful for forming a proteinaceous gel which may then be used for healing wounds, or in
                                                                                                                                                                                                                                                                             1723 erdrkfreeeglrggreegglrsg-esdrkfreeeglrgereegglrpggrdgkyrwee 1780
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human
wound
                         Homo sapiens
                                                                                                                                      Human unliganded androgen receptor protein.
                                                                                                                                                                19-JAN-2000 (first entry)
                                                                                                                                                                                                                 Y33504 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Fig 3A-W; 126pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1999-561041/47
N-PSDB; Z22301.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kim I,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US5958752-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             breast implant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; trichohyalin; TRHY;
terminally differentiating
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A human trichohyalin (TRHY) protein
                                                                                                                                                                                                                                                                                                                                                                                                                       breast implants
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   30-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           14-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    28-SEP-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (USSH ) US DEPT HEALTH & HUMAN SERVICES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-NOV-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Y30795 standard; Protein; 1898
                                                                                                                                                                                                                                          14
                                                                                                                                                                                                                                                                                            \ensuremath{\operatorname{trichohyalin}} useful for forming a proteinaceous gel that promotes healing -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chung S,
                                                                                                                                                                                                                                                                                                                                                                                                1898 AA;
                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   93US-0056200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           97US-0800644
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Park S,
                                                                                                                                                                                                                                                                                                                                              20.3%;
                                                                                                                                                                                                                    154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        protein; tissue structure; wound healing;
epidermal tissue; proteinaceous gel;
                                                                                                                                                                                                                                                                                                                                  10;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Steinert PM,
                                                                                                                                                                                                                                                                                                                               Score 71.5; DB
Pred. No. 5.1;
10; Mismatches
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                                                                                                                                                                                                                                                                                                                                                          DB 20;
                                                                                                                                                                                                                                                                                                                                 24;
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                                                                                                                                                                                                                                                                                                                                                         Length 1898;
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16-SEP-1999

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RESULT 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention describes novel pure proapoptotic dependence peptides which comprise a sequence of an active dependence domain selected from dependence polypeptides consisting of p75NTR, androgen receptor, DCC, huntingtin polypeptide, Machado-Joseph disease gene product, SCA1, SCA2, SCA6 and atrophin-1 polypeptide. The proapoptotic peptides are capable of inducing cell death and can be used to develop products to mediate or inhibit apoptosis. The methods can be used for reducing the severity of a proapoptotic dependence domain mediated pathological conditions e.g. Huntington's disease, Alzheimer's disease, Kennedy's disease, Spinocerebellar ataxias, dentatorubropallidoluysian atrophy, Machado-Joseph disease, stroke or head trauma. They can also be used for mediated pathological conditions be used for several traumants.
WPI; 1998-080076/08
                            (KISH/) KISHIMOTO C.
(SEKI ) SEKISUI CHEM IND CO LTD.
                                                                          27-MAY-1996;
                                                                                                           27-MAY-1996;
                                                                                                                                           09-DEC-1997
                                                                                                                                                                          JP09313185-A
                                                                                                                                                                                                        Mus sp
                                                                                                                                                                                                                                       Mouse; human;
hHDGF; HET-A;
                                                                                                                                                                                                                                                                                 Mouse liver cancer-originated culture cell growth factor
                                                                                                                                                                                                                                                                                                                   27-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                               W37483 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    reducing the severity of a pathological condition mediated by upregulated cell proliferation or cell survival e.g. neoplastic, malignant, autoimmune or fibrotic conditions. This sequence represents a human unliganded androgen receptor described in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New proapoptotic dependence peptides, used to develop products treating, e.g. Alzheimer's disease -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Page 178-179; 199pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1999-561617/47.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        58 qqqqqq-----qqqqqqqqqqqqetsprqqqqqqgedgspqahrrgptgylvldeeqq 109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5 QQREYEDCRRRCEQQEPRQQYQCQRRCREQQRQHGRGGDLINPQRGGSGRYEEGEEKQ 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        154 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                                                                                       HET-B
                                                                                                                                                                                                                                                   liver cancer-originated culture cell growth factor;
                                                                            96JP-0131788
                                                                                                           96JP-0131788
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                                                                                                                                                                                                                                                                                                                                                                                 669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Score 71; DB 20; Pred. No. 0.45; 12; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 20; Length 154;
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                                                                                                                                               Query Match
Best Local Similarity 35.0
                                                                                                                                                                               polypeptide protein similar to protein of (1), but comprises no protein combined naturally and also has 2 amino acid sequences (as given in the specification) which may be modified to produce a sequence of at least 5 amino acids; (3) a recombinant DNA molecule comprising a vector and the DNA segment of (1); (4) a cell comprising (3); (5) producing a polypeptide comprising amino acids corresponding to a protein of (1), in which a cell comprising (3) is cultured under a conditions enabling the expression of (1) to produce the polypeptide which is isolated. The above method may be used to produce proteins HET-A and HET-B.
                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                      The present sequence represents a protein which has local homology to human liver cancer-originated culture cell isolated from a mouse testLole cDNA library. The sequence may (1) be modified to produce a polypeptide comprising at least 5 continuous amino acids; (2) a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 10; Page 13-14; 18pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DNA segment encoding protein homologous to cancer-originated culture cell - which may polypeptide of at least 5 continuous amino
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            N-PSDB; V01731
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       producing, e.g. protein HET-A
1 KRDPQQREYEDCRRRCEQQEPRQQYQCQRRCREQQRQHGRGGDLINPQRGGSGRYEEGEE 60
                                                                                                                                               669 AA;
                                                              20.1%;
                                      Score 71; DB Pred. No. 2; 8; Mismatches
                                             8;
                                                                                 DB 19; Length 669;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        human liver
be modified to produce
acids, useful in
                                             25;
                                          Indels
                                          6;
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